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<110> Li, En Okano, Masaki Xie, Shaoping Chen, Taiping

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ctgtaccctg ccattcccgc agcccgaagg cggcccattc gagtcctgtc attgtttgat 1860 ggcatcgcga caggctacct agtcctcaaa gagttgggca taaaggtagg aaagtacgtc 1920 gcttctgaag tgtgtgagga gtccattgct gttggaaccg tgaagcacga ggggaatatc 1980 aaatacgtga acgacgtgag gaacatcaca aagaaaaata ttgaagaatg gggcccattt 2040 gacttggtga ttggcggaag cccatgcaac gatctctcaa atgtgaatcc agccaggaaa 2100 ggcctgtatg agggtacagg ccggctcttc ttcgaatttt accacctgct gaattactca 2160 cgccccaagg agggtgatga ccggccgttc ttctggatgt ttgagaatgt tgtagccatg 2220 aaggttggcg acaagaggga catctcacgg ttcctggagt gtaatccagt gatgattgat 2280 gccatcaaag tttctgctgc tcacagggcc cgatacttct ggggcaacct acccgggatg 2340 aacaggcccg tgatagcatc aaagaatgat aaactcgagc tgcaggactg cttggaatac 2400 aataggatag ccaagttaaa gaaagtacag acaataacca ccaagtcgaa ctcgatcaaa 2460 caggggaaaa accaactttt ccctgttgtc atgaatggca aagaagatgt tttgtggtgc 2520 actgageteg aaaggatett tggettteet gtgeactaea cagaegtgte caacatggge 2580 cgtggtgccc gccagaagct gctgggaagg tcctggagcg tgcctgtcat ccgacacctc 2640 ttcgcccctc tgaaggacta ctttgcatgt gaatagttcc agccaggccc caagcccact 2700 ggggtgtgtg gcagagccag gaccaggag gtgtgattcc tgaaggcatc cccaggccct 2760 gctcttcctc agctgtgtgg gtcataccgt gtacctcagt tccctcttgc tcagtggggg 2820 cagagecace tgaetettge aggggtagee tgaggtgeeg ceteettgtg cacaaateag 2880 acctggctgc ttggagcagc ctaacacggt gctcattttt tcttctccta aaactttaaa 2940 acttgaagta ggtagcaacg tggctttttt tttttccctt cctgggtcta ccactcagag 3000 aaacaatggc taagatacca aaaccacagt gccgacagct ctccaatact caggttaatg 3060 ctgaaaaatc atccaagaca gttattgcaa gagtttaatt tttgaaaact gggtactgct 3120 atgtgtttac agacgtgtgc agttgtaggc atgtagctac aggacatttt taagggccca 3180 ggatcgtttt ttcccagggc aagcagaaga gaaaatgttg tatatgtctt ttacccggca 3240 cattcccctt gcctaaatac aagggctgga gtctgcacgg gacctattag agtattttcc 3300 acaatgatga tgatttcagc agggatgacg tcatcatcac attcagggct atttttccc 3360 ccacaaaccc aagggcaggg gccactctta gctaaatccc tccccgtgac tgcaatagaa 3420 ccctctgggg agctcaggaa ggggtgtgct gagttctata atataagctg ccatatattt 3480 tgtagacaag tatggctcct ccatatctcc ctcttcccta ggagaggagt gtgaagcaag 3540 gagettagat aagacaceee etcaaaceea tteeetetee aggagaceta eeetecacag 3600 gcacaggtcc ccagatgaga agtctgctac cctcatttct catctttta ctaaactcag 3660 aggcagtgac agcagtcagg gacagacata catttctcat accttcccca catctgagag 3720 atgacaggga aaactgcaaa gctcggtgct ccctttggag attttttaat ccttttttat 3780 tccataagaa gtcgttttta gggagaacgg gaattcagac aagctgcatt tcagaaatgc 3840 tgtcataatg gtttttaaca ccttttactc ttcttactgg tgctattttg tagaataagg 3900 aacaacgttg acaagttttg tggggctttt tatacacttt ttaaaaatctc aaacttctat 3960 ttttatgttt aacgttttca ttaaaatttt tttgtaactg gagccacgac gtaacaaata 4020 tggggaaaaa actgtgcctt gtttcaacag tttttgctaa tttttaggct gaaagatgac 4080 ggatgcctag agtttacctt atgtttaatt aaaatcagta tttgtctaaa aaaaaaaaa 4140 aaaaa 4145

<212> PRT

<213> Mus musculus

<400> 5

Met Pro Ser Ser Gly Pro Gly Asp Thr Ser Ser Ser Ser Leu Glu Arg

1 5 10 15

Glu Asp Asp Arg Lys Glu Gly Glu Glu Glu Glu Glu Asn Arg Gly Lys
20 25 30

Glu Glu Arg Gln Glu Pro Ser Ala Thr Ala Arg Lys Val Gly Arg Pro 35 40 45

Gly Arg Lys Arg Lys His Pro Pro Val Glu Ser Ser Asp Thr Pro Lys
50 55 60

Asp Pro Ala Val Thr Thr Lys Ser Gln Pro Met Ala Gln Asp Ser Gly 65 70 75 80

Pro Ser Asp Leu Leu Pro Asn Gly Asp Leu Glu Lys Arg Ser Glu Pro 85 90 95

Gln Pro Glu Glu Gly Ser Pro Ala Ala Gly Gln Lys Gly Gly Ala Pro 100 105 110

Ala Glu Gly Glu Gly Thr Glu Thr Pro Pro Glu Ala Ser Arg Ala Val 115 120 125

Glu Asn Gly Cys Cys Val Thr Lys Glu Gly Arg Gly Ala Ser Ala Gly 130 135 140

Gly Ser Arg Gly Arg Leu Arg Gly Gly Leu Gly Trp Glu Ser Ser Leu 165 170 175

Arg Gln Arg Pro Met Pro Arg Leu Thr Phe Gln Ala Gly Asp Pro Tyr
180 185 190

Tyr Ile Ser Lys Arg Lys Arg Asp Glu Trp Leu Ala Arg Trp Lys Arg 195 200 205

- Glu Ala Glu Lys Lys Ala Lys Val Ile Ala Val Met Asn Ala Val Glu 210 215 220
- Glu Asn Gln Ala Ser Gly Glu Ser Gln Lys Val Glu Glu Ala Ser Pro 225 230 235 240
- Pro Ala Val Gln Gln Pro Thr Asp Pro Ala Ser Pro Thr Val Ala Thr
 245 250 255
- Thr Pro Glu Pro Val Gly Gly Asp Ala Gly Asp Lys Asn Ala Thr Lys
 260 265 270
- Ala Ala Asp Asp Glu Pro Glu Tyr Glu Asp Gly Arg Gly Phe Gly Ile 275 280 285
- Gly Glu Leu Val Trp Gly Lys Leu Arg Gly Phe Ser Trp Trp Pro Gly
 290 295 300
- Arg Ile Val Ser Trp Trp Met Thr Gly Arg Ser Arg Ala Ala Glu Gly 305 310 315 320
- Thr Arg Trp Val Met Trp Phe Gly Asp Gly Lys Phe Ser Val Val Cys 325 330 335
- Val Glu Lys Leu Met Pro Leu Ser Ser Phe Cys Ser Ala Phe His Gln 340 345 350
- Ala Thr Tyr Asn Lys Gln Pro Met Tyr Arg Lys Ala Ile Tyr Glu Val ... 355 360 365
- Leu Gln Val Ala Ser Ser Arg Ala Gly Lys Leu Phe Pro Ala Cys His 370 375 380
- Asp Ser Asp Glu Ser Asp Ser Gly Lys Ala Val Glu Val Gln Asn Lys 385 390 395 400
- Gln Met Ile Glu Trp Ala Leu Gly Gly Phe Gln Pro Ser Gly Pro Lys 405 410 415
- Gly Leu Glu Pro Pro Glu Glu Glu Lys Asn Pro Tyr Lys Glu Val Tyr
 420 425 430
- Thr Asp Met Trp Val Glu Pro Glu Ala Ala Ala Tyr Ala Pro Pro

Pro Ala Lys Lys Pro Arg Lys Ser Thr Thr Glu Lys Pro Lys Val Lys Glu Ile Ile Asp Glu Arg Thr Arg Glu Arg Leu Val Tyr Glu Val Arg Gln Lys Cys Arg Asn Ile Glu Asp Ile Cys Ile Ser Cys Gly Ser Leu Asn Val Thr Leu Glu His Pro Leu Phe Ile Gly Gly Met Cys Gln Asn Cys Lys Asn Cys Phe Leu Glu Cys Ala Tyr Gln Tyr Asp Asp Asp Gly Tyr Gln Ser Tyr Cys Thr Ile Cys Cys Gly Gly Arg Glu Val Leu Met Cys Gly Asn Asn Asn Cys Cys Arg Cys Phe Cys Val Glu Cys Val Asp [′]555 Leu Leu Val Gly Pro Gly Ala Ala Gln Ala Ala Ile Lys Glu Asp Pro Trp Asn Cys Tyr Met Cys Gly His Lys Gly Thr Tyr Gly Leu Leu Arg Arg Arg Glu Asp Trp Pro Ser Arg Leu Gln Met Phe Phe Ala Asn Asn

Thr Gly Leu Leu Val Leu Lys Asp Leu Gly Ile Gln Val Asp Arg Tyr 645 650 655

Ile Ala Ser Glu Val Cys Glu Asp Ser Ile Thr Val Gly Met Val Arg 660 665 670

His Asp Gln Glu Phe Asp Pro Pro Lys Val Tyr Pro Pro Val Pro Ala

Glu Lys Arg Lys Pro Ile Arg Val Leu Ser Leu Phe Asp Gly Ile Ala

610 .

- His Gln Gly Lys Ile Met Tyr Val Gly Asp Val Arg Ser Val Thr Gln 675 680 685
- Lys His Ile Gln Glu Trp Gly Pro Phe Asp Leu Val Ile Gly Gly Ser 690 695 700
- Pro Cys Asn Asp Leu Ser Ile Val Asn Pro Ala Arg Lys Gly Leu Tyr 705 710 715 720
- Glu Gly Thr Gly Arg Leu Phe Phe Glu Phe Tyr Arg Leu Leu His Asp
 725 730 735
- Ala Arg Pro Lys Glu Gly Asp Asp Arg Pro Phe Phe Trp Leu Phe Glu
 740 745 750
- Asn Val Val Ala Met Gly Val Ser Asp Lys Arg Asp Ile Ser Arg Phe 755 760 765
- Leu Glu Ser Asn Pro Val Met Ile Asp Ala Lys Glu Val Ser Ala Ala 770 775 780
- His Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro Gly Met Asn Arg Pro 785 790 795 800
- Leu Ala Ser Thr Val Asn Asp Lys Leu Glu Leu Gln Glu Cys Leu Glu 805 810 815
- His Gly Arg Ile Ala Lys Phe Ser Lys Val Arg Thr Ile Thr Thr Arg 820 825 830
- Ser Asn Ser Ile Lys Gln Gly Lys Asp Gln His Phe Pro Val Phe Met 835 840 845
- Asn Glu Lys Glu Asp Ile Leu Trp Cys Thr Glu Met Glu Arg Val Phe 850 855 860
- Gly Phe Pro Val His Tyr Thr Asp Val Ser Asn Met Ser Arg Leu Ala 865 870 875 880
- Arg Gln Arg Leu Leu Gly Arg Ser Trp Ser Val Pro Val Ile Arg His 885 890 895

Leu Phe Ala Pro Leu Lys Glu Tyr Phe Ala Cys Val 900 905

<210> 6

<211> 859

<212> PRT

<213> Mus musculus

<400> 6

Met Lys Gly Asp Ser Arg His Leu Asn Glu Glu Glu Gly Ala Ser Gly

1 5 10 15

Tyr Glu Glu Cys Ile Ile Val Asn Gly Asn Phe Ser Asp Gln Ser Ser
20 25 30

Asp Thr Lys Asp Ala Pro Ser Pro Pro Val Leu Glu Ala Ile Cys Thr 35 40 45

Glu Pro Val Cys Thr Pro Glu Thr Arg Gly Arg Arg Ser Ser Arg
50 55 60

Leu Ser Lys Arg Glu Val Ser Ser Leu Leu Asn Tyr Thr Gln Asp Met 65 70 75 80

Thr Gly Asp Gly Asp Arg Asp Glu Val Asp Asp Gly Asn Gly Ser 85 90 95

Asp Ile Leu Met Pro Lys Leu Thr Arg Glu Thr Lys Asp Thr Arg Thr
100 105 110

Arg Ser Glu Ser Pro Ala Val Arg Thr Arg His Ser Asn Gly Thr Ser
115 120 125

Ser Leu Glu Arg Gln Arg Ala Ser Pro Arg Ile Thr Arg Gly Arg Gln 130 135 140

Gly Arg His His Val Gln Glu Tyr Pro Val Glu Phe Pro Ala Thr Arg 145 150 155 160

Ser Arg Arg Arg Ala Ser Ser Ser Ala Ser Thr Pro Trp Ser Ser 165 170 175

- Pro Ala Ser Val Asp Phe Met Glu Glu Val Thr Pro Lys Ser Val Ser 180 185 190
- Thr Pro Ser Val Asp Leu Ser Gln Asp Gly Asp Gln Glu Gly Met Asp 195 200 205
- Thr Thr Gln Val Asp Ala Glu Ser Arg Asp Gly Asp Ser Thr Glu Tyr 210 215 220
- Gln Asp Asp Lys Glu Phe Gly Ile Gly Asp Leu Val Trp Gly Lys Ile 225 230 235 240
- Lys Gly Phe Ser Trp Trp Pro Ala Met Val Val Ser Trp Lys Ala Thr
 245 250 255
- Ser Lys Arg Gln Ala Met Pro Gly Met Arg Trp Val Gln Trp Phe Gly
 260 265 270
- Asp Gly Lys Phe Ser Glu Ile Ser Ala Asp Lys Leu Val Ala Leu Gly 275 280 285
- Leu Phe Ser Gln His Phe Asn Leu Ala Thr Phe Asn Lys Leu Val Ser 290 295 300
- Tyr Arg Lys Ala Met Tyr His Thr Leu Glu Lys Ala Arg Val Arg Ala 305 310 315 320
- Gly Lys Thr Phe Ser Ser Pro Gly Glu Ser Leu Glu Asp Gln Leu
 325 330 335
- Lys Pro Met Leu Glu Trp Ala His Gly Gly Phe Lys Pro Thr Gly Ile 340 345 350
- Glu Gly Leu Lys Pro Asn Lys Lys Gln Pro Val Val Asn Lys Ser Lys 355 360 365
- Val Arg Arg Ser Asp Ser Arg Asn Leu Glu Pro Arg Arg Arg Glu Asn 370 380
- Lys Ser Arg Arg Arg Thr Thr Asn Asp Ser Ala Ala Ser Glu Ser Pro 385 390 395 400
- Pro Pro Lys Arg Leu Lys Thr Asn Ser Tyr Gly Gly Lys Asp Arg Gly

Glu Asp Glu Glu Ser Arg Glu Arg Met Ala Ser Glu Val Thr Asn Asn Lys Gly Asn Leu Glu Asp Arg Cys Leu Ser Cys Gly Lys Lys Asn Pro Val Ser Phe His Pro Leu Phe Glu Gly Gly Leu Cys Gln Ser Cys Arg Asp Arg Phe Leu Glu Leu Phe Tyr Met Tyr Asp Glu Asp Gly Tyr Gln Ser Tyr Cys Thr Val Cys Cys Glu Gly Arg Glu Leu Leu Cys Ser Asn Thr Ser Cys Cys Arg Cys Phe Cys Val Glu Cys Leu Glu Val Leu Val Gly Ala Gly Thr Ala Glu Asp Ala Lys Leu Gln Glu Pro Trp Ser Cys Tyr Met Cys Leu Pro Gln Arg Cys His Gly Val Leu Arg Arg Arg Lys Asp Trp Asn Met Arg Leu Gln Asp Phe Phe Thr Thr Asp Pro Asp Leu Glu Glu Phe Glu Pro Pro Lys Leu Tyr Pro Ala Ile Pro Ala Ala Lys Arg Arg Pro Ile Arg Val Leu Ser Leu Phe Asp Gly Ile Ala Thr Gly Tyr Leu Val Leu Lys Glu Leu Gly Ile Lys Val Glu Lys Tyr Ile Ala Ser Glu Val Cys Ala Glu Ser Ile Ala Val Gly Thr Val Lys His Glu Gly Gln Ile Lys Tyr Val Asn Asp Val Arg Lys Ile Thr Lys Lys

- Asn Ile Glu Glu Trp Gly Pro Phe Asp Leu Val Ile Gly Gly Ser Pro 645 650 655
- Cys Asn Asp Leu Ser Asn Val Asn Pro Ala Arg Lys Gly Leu Tyr Glu 660 665 670
- Gly Thr Gly Arg Leu Phe Phe Glu Phe Tyr His Leu Leu Asn Tyr Thr 675 680 685
- Arg Pro Lys Glu Gly Asp Asn Arg Pro Phe Phe Trp Met Phe Glu Asn 690 695 700
- Val Val Ala Met Lys Val Asn Asp Lys Lys Asp Ile Ser Arg Phe Leu 705 710 715 720
- Ala Cys Asn Pro Val Met Ile Asp Ala Ile Lys Val Ser Ala Ala His
 725 730 735
- Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro Gly Met Asn Arg Pro Val
 740 745 750
- Met Ala Ser Lys Asn Asp Lys Leu Glu Leu Gln Asp Cys Leu Glu Phe
 755 760 765
- Ser Arg Thr Ala Lys Leu Lys Lys Val Gln Thr Ile Thr Thr Lys Ser 770 780
- Asn Ser Ile Arg Gln Gly Lys Asn Gln Leu Phe Pro Val Val Met Asn 785 790 795 800
- Gly Lys Asp Asp Val Leu Trp Cys Thr Glu Leu Glu Arg Ile Phe Gly 805 810 815
- Phe Pro Ala His Tyr Thr Asp Val Ser Asn Met Gly Arg Gly Ala Arg 820 825 830 $^{\circ}$
- Gln Lys Leu Leu Gly Arg Ser Trp Ser Val Pro Val Ile Arg His Leu 835 840 845
- Phe Ala Pro Leu Lys Asp Tyr Phe Ala Cys Glu 850 855

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<211> 912

<212> PRT

<213> Homo sapiens

<400> 7

Met Pro Ala Met Pro Ser Ser Gly Pro Gly Asp Thr Ser Ser Ala
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Ala Glu Arg Glu Glu Asp Arg Lys Asp Gly Glu Glu Glu Glu Pro 20 25 30

Arg Gly Lys Glu Glu Arg Gln Glu Pro Ser Thr Thr Ala Arg Lys Val
35 40 45

Gly Arg Pro Gly Arg Lys Arg Lys His Pro Pro Val Glu Ser Gly Asp
50 55 60

Thr Pro Lys Asp Pro Ala Val Ile Ser Lys Ser Pro Ser Met Ala Gln 65 70 75 80

Asp Ser Gly Ala Ser Glu Leu Leu Pro Asn Gly Asp Leu Glu Lys Arg 85 90 95

Ser Glu Pro Gln Pro Glu Glu Gly Ser Pro Ala Gly Gln Lys Gly
100 105 110

Gly Ala Pro Ala Glu Gly Glu Gly Ala Ala Glu Thr Leu Pro Glu Ala 115 120 125

Ser Arg Ala Val Glu Asn Gly Cys Cys Thr Pro Lys Glu Gly Arg Gly 130 135 140

Met Lys Met Glu Gly Ser Arg Gly Arg Leu Arg Gly Gly Leu Gly Trp 165 170 175

Glu Ser Ser Leu Arg Gln Arg Pro Met Pro Arg Leu Thr Phe Gln Ala 180 185 190

- Gly Asp Pro Tyr Tyr Ile Ser Lys Arg Lys Arg Asp Glu Trp Leu Ala 195 200 205
- Arg Trp Lys Arg Glu Ala Glu Lys Lys Ala Lys Val Ile Ala Gly Met 210 215 220
- Asn Ala Val Glu Glu Asn Gln Gly Pro Gly Glu Ser Gln Lys Val Glu 225 230 235 240
- Glu Ala Ser Pro Pro Ala Val Gln Gln Pro Thr Asp Pro Ala Ser Pro 245 250 255
- Thr Val Ala Thr Thr Pro Glu Pro Val Gly Ser Asp Ala Gly Asp Lys
 260 265 270
- Asn Ala Thr Lys Ala Gly Asp Asp Glu Pro Glu Tyr Glu Asp Gly Arg 275 280 285
- Gly Phe Gly Ile Gly Glu Leu Val Trp Gly Lys Leu Arg Gly Phe Ser 290 295 300
- Trp Trp Pro Gly Arg Ile Val Ser Trp Trp Met Thr Gly Arg Ser Arg 305 310 315 320
- Ala Ala Glu Gly Thr Arg Trp Val Met Trp Phe Gly Asp Gly Lys Phe 325 330 335
- Ser Val Val Cys Val Glu Lys Leu Met Pro Leu Ser Ser Phe Cys Ser 340 345 , 350
- Ala Phe His Gln Ala Thr Tyr Asn Lys Gln Pro Met Tyr Arg Lys Ala 355 360 365
- Ile Tyr Glu Val Leu Gln Val Ala Ser Ser Arg Ala Gly Lys Leu Phe 370 375 380
- Pro Val Cys His Asp Ser Asp Glu Ser Asp Thr Ala Lys Ala Val Glu 385 390 395 400
- Val Gln Asn Lys Pro Met Ile Glu Trp Ala Leu Gly Gly Phe Gln Pro 405 410 415
- Ser Gly Pro Lys Gly Leu Glu Pro Pro Glu Glu Lys Asn Pro Tyr

Lys Glu Val Tyr Thr Asp Met Trp Val Glu Pro Glu Ala Ala Ala Tyr Ala Pro Pro Pro Pro Ala Lys Lys Pro Arg Lys Ser Thr Ala Glu Lys Pro Lys Val Lys Glu Ile Ile Asp Glu Arg Thr Arg Glu Arg Leu Val Tyr Glu Val Arg Gln Lys Cys Arg Asn Ile Glu Asp Ile Cys Ile Ser Cys Gly Ser Leu Asn Val Thr Leu Glu His Pro Leu Phe Val Gly Gly Met Cys Gln Asn Cys Lys Asn Cys Phe Leu Glu Cys Ala Tyr Gln Tyr 525、 Asp Asp Gly Tyr Gln Ser Tyr Cys Thr Ile Cys Cys Gly Gly Arg Glu Val Leu Met Cys Gly Asn Asn Cys Cys Arg Cys Phe Cys Val Glu Cys Val Asp Leu Leu Val Gly Pro Gly Ala Ala Gln Ala Ala Ile Lys Glu Asp Pro Trp Asn Cys Tyr Met Cys Gly His Lys Gly Thr Tyr Gly Leu Leu Arg Arg Glu Asp Trp Pro Ser Arg Leu Gln Met Phe Phe Ala Asn Asn His Asp Gln Glu Phe Asp Pro Pro Lys Val Tyr Pro Pro Val Pro Ala Glu Lys Arg Lys Pro Ile Arg Val Leu Ser Leu Phe Asp Gly Ile Ala Thr Gly Leu Leu Val Leu Lys Asp Leu Gly Ile Gln

- Val Asp Arg Tyr Ile Ala Ser Glu Val Cys Glu Asp Ser Ile Thr Val 660 665 670
- Gly Met Val Arg His Gln Gly Lys Ile Met Tyr Val Gly Asp Val Arg
 675 680 685
- Ser Val Thr Gln Lys His Ile Gln Glu Trp Gly Pro Phe Asp Leu Val 690 695 700
- Ile Gly Gly Ser Pro Cys Asn Asp Leu Ser Ile Val Asn Pro Ala Arg
 705 710 715 720
- Lys Gly Leu Tyr Glu Gly Thr Gly Arg Leu Phe Phe Glu Phe Tyr Arg
 725 730 735
- Leu Leu His Asp Ala Arg Pro Lys Glu Gly Asp Asp Arg Pro Phe Phe 740 745 750
- Trp Leu Phe Glu Asn Val Val Ala Met Gly Val Ser Asp Lys Arg Asp 755 760 765
- Ile Ser Arg Phe Leu Glu Ser Asn Pro Val Met Ile Asp Ala Lys Glu
 770 780
- Val Ser Ala Ala His Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro Gly 785 790 795 800
- Met Asn Arg Pro Leu Ala Ser Thr Val Asn Asp Lys Leu Glu Leu Gln 805 810 815
- Glu Cys Leu Glu His Gly Arg Ile Ala Lys Phe Ser Lys Val Arg Thr 820 825 830
- Ile Thr Thr Arg Ser Asn Ser Ile Lys Gln Gly Lys Asp Gln His Phe 835 840 845
- Pro Val Phe Met Asn Glu Lys Glu Asp Ile Leu Trp Cys Thr Glu Met 850 855 860
- Glu Arg Val Phe Gly Phe Pro Val His Tyr Thr Asp Val Ser Asn Met 865 870 875 880

Ser Arg Leu Ala Arg Gln Arg Leu Leu Gly Arg Ser Trp Ser Val Pro 885 890 895

Val Ile Arg His Leu Phe Ala Pro Leu Lys Glu Tyr Phe Ala Cys Val 900 905 910

<210> 8

<211> 853

<212> PRT

<213> Homo sapiens

<400> 8

Met Lys Gly Asp Thr Arg His Leu Asn Gly Glu Glu Asp Ala Gly Gly

1 10 15

Arg Glu Asp Ser Ile Leu Val Asn Gly Ala Cys Ser Asp Gln Ser Ser 20 25 30

Asp Ser Pro Pro Ile Leu Glu Ala Ile Arg Thr Pro Glu Ile Arg Gly 35 40 45 .

Arg Arg Ser Ser Ser Arg Leu Ser Lys Arg Glu Val Ser Ser Leu Leu 50 55 60

Ser Tyr Thr Gln Asp Leu Thr Gly Asp Gly Asp Gly Glu Asp Gly Asp 65 70 75 80

Gly Ser Asp Thr Pro Val Met Pro Lys Leu Phe Arg Glu Thr Arg Thr
85 90 95

Arg Ser Glu Ser Pro Ala Val Arg Thr Arg Asn Asn Asn Ser Val Ser
100 105 110

Ser Arg Glu Arg His Arg Pro Ser Pro Arg Ser Thr Arg Gly Arg Gln
115 120 125

Gly Arg Asn His Val Asp Glu Ser Pro Val Glu Phe Pro Ala Thr Arg 130 135 140

- Ser Leu Arg Arg Arg Ala Thr Ala Ser Ala Gly Thr Pro Trp Pro Ser 145 150 155 160
- Pro Pro Ser Ser Tyr Leu Thr Ile Asp Leu Thr Asp Asp Thr Glu Asp 165 170 175
- Thr His Gly Thr Pro Gln Ser Ser Ser Thr Pro Tyr Ala Arg Leu Ala 180 185 190
- Gln Asp Ser Gln Gln Gly Gly Met Glu Ser Pro Gln Val Glu Ala Asp 195 200 205
- Ser Gly Asp Gly Asp Ser Ser Glu Tyr Gln Asp Gly Lys Glu Phe Gly 210 215 220
- Ile Gly Asp Leu Val Trp Gly Lys Ile Lys Gly Phe Ser Trp Trp Pro 225 230 235 240
- Ala Met Val Val Ser Trp Lys Ala Thr Ser Lys Arg Gln Ala Met Ser 245 250 255
- Gly Met Arg Trp Val Gln Trp Phe Gly Asp Gly Lys Phe Ser Glu Val 260 265 270
- Ser Ala Asp Lys Leu Val Ala Leu Gly Leu Phe Ser Gln His Phe Asn 275 . 280 285
- Leu Ala Thr Phe Asn Lys Leu Val Ser Tyr Arg Lys Ala Met Tyr His 290 295 300
- Ala Leu Glu Lys Ala Arg Val Arg Ala Gly Lys Thr Phe Pro Ser Ser 305 310 315 320
- Pro Gly Asp Ser Leu Glu Asp Gln Leu Lys Pro Met Leu Glu Trp Ala 325 330 335
- His Gly Gly Phe Lys Pro Thr Gly Ile Glu Gly Leu Lys Pro Asn Asn 340 345 350
- Thr Gln Pro Val Val Asn Lys Ser Lys Val Arg Arg Ala Gly Ser Arg 355 360 365
- Lys Leu Glu Ser Arg Lys Tyr Glu Asn Lys Thr Arg Arg Arg Thr Ala

	370					375					380				
Asp 385	Asp	Ser	Ala	Thr	Ser 390	Asp	Tyr	Cys	Pro	Ala 395	Pro	Lys	Arg	Leu	Lys 400
Thr	Asn	Cys	Tyr	Asn 405	Asn	Gly	Lys	Asp	Arg 410	Gly	Asp	Glu	Asp	Gln 415	Ser
Arg	Glu	Gln	Met 420	Ala	Ser	Asp	Val	Ala 425	Asn	Asn	Lys	Ser	Ser 430	Leu	Glu
Asp	Gly	Cys 435	Leu	Ser	Сув	Gly	Arg 440	Lys	Asn	Pro	Val	Ser 445	Phe	His	Pro
Leu	Phe 450	Glu	Gly	Gly	Leu	Cys 455	Gln	Thr	Cys	Arg	Asp 460	Arg	Phe	Leu	Glu
Leu 465	Phe	Tyr	Met	Tyr	Asp 470	Asp	Asp	Gly	Tyr	Gln 475	Ser	Tyr	Cys	Thr	Val
Сув	Cys	Glu	Gly	Arg 485	Glu	Leu	Leu	Leu	Cys 490	Ser	Asn	Thr	Ser	Cys 495	Cys
Arg	Cys	Phe	Cys 500	Val	Glu	Cys	Leu	Glu 505	Val •	Leu	Val	Gly	Thr 510	Gly	Thr
Ala	Ala	Glu 515	Ala	Lys	Leu	Gln	Glu 520	Pro	Trp	Ser	Cys	Tyr 525	Met	Cys	Lev
Pro	Gln 530	Arg	Cys	His	Gly	Val 535	Leu	Arg	Arg	Arg	Lys 540	Asp	Trp	Asn	Val
Arg 545	Leu	Gln	Ala	Phe	Phe 550	Thr	Ser	Asp	Thr	Gly 555	Leu	Glu	Tyr	Glu	Ala 560
Pro	Lys	Leu	Tyr	Pro 565	Ala	Ile	Pro	Ala	Ala 570	Arg	Arg	Arg	Pro	Ile 575	Arg
Val	Leu	Ser	Leu 580	Phe	Asp	Gly	Ile	Ala 585	Thr	Gly	Tyr	Leu	Val 590	Leu	Lys

Glu Leu Gly Ile Lys Val Gly Lys Tyr Val Ala Ser Glu Val Cys Glu

605

600

595

- Glu Ser Ile Ala Val Gly Thr Val Lys His Glu Gly Asn Ile Lys Tyr 610 615 620
- Val Asn Asp Val Arg Asn Ile Thr Lys Lys Asn Ile Glu Glu Trp Gly 625 630 635 640
- Pro Phe Asp Leu Val Ile Gly Gly Ser Pro Cys Asn Asp Leu Ser Asn 645 650 655
- Val Asn Pro Ala Arg Lys Gly Leu Tyr Glu Gly Thr Gly Arg Leu Phe 660 665 670
- Phe Glu Phe Tyr His Leu Leu Asn Tyr Ser Arg Pro Lys Glu Gly Asp 675 680 685
- Asp Arg Pro Phe Phe Trp Met Phe Glu Asn Val Val Ala Met Lys Val 690 695 700
- Gly Asp Lys Arg Asp Ile Ser Arg Phe Leu Glu Cys Asn Pro Val Met 705 710 715 720
- Ile Asp Ala Ile Lys Val Ser Ala Ala His Arg Ala Arg Tyr Phe Trp
 725 730 735
- Gly Asn Leu Pro Gly Met Asn Arg Pro Val Ile Ala Ser Lys Asn Asp
 740 745 750
- Lys Leu Glu Leu Gln Asp Cys Leu Glu Tyr Asn Arg Ile Ala Lys Leu 755 760 765
- Lys Lys Val Gln Thr Ile Thr Thr Lys Ser Asn Ser Ile Lys Gln Gly 770 775 780
- Lys Asn Gln Leu Phe Pro Val Val Met Asn Gly Lys Glu Asp Val Leu 785 790 795 800
- Trp Cys Thr Glu Leu Glu Arg Ile Phe Gly Phe Pro Val His Tyr Thr 805 810 815
- Asp Val Ser Asn Met Gly Arg Gly Ala Arg Gln Lys Leu Leu Gly Arg 820 825 830

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cggggctccc ctgcaaggcc aggaggctct gctcccactg agcaggagaa agctgaggta 240
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<400> 33

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<212> DNA

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aagggttatt gcatgagtct ggatgaatcc cactctcagc tgtccacggg cccgaccacc 240
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cacgcaaaac agaacccagt tagcagcagg gagacgagaa caccacacaa gacatttttc 180
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cgctgttacc tcttgtttac agtttatata tatatgatag atatgagatn tatatataaa 300
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geggeacatt tgeceteeca gecaetgage tgtegtgeea geaecattee tgggteaege 240
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 aaagggttat tgcatgagtc tggatgaatc ccactctcag ctgtccacgg gcccgaccac 240
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 ctttccattt cagtgcacca taagatgtcc tctttctcat tcatgaagac aggaaaaatg 240
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aggagggaga tgatcgcccc ttctctggct ctttgagaat ttggtggcca tggcgttagt 240
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agaagtetea tetgeacaga ggeeeeteta ettetggggt caceteeeeg tattaacagg 360
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ncatgeteca gacaetteeg nagggteaca acagaageat nttecagggg gtggnggeea 480
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tagatatgag atatatatat aaaaggtact gttaactact gtacaacccg acttcataat 180
ggtgctttca aacagcgaga tgagtaaaaa catcagcttc cacgttgcct tctgcgcaaa 240
gggtttcacc aaggatggag aaagggagac agcttgcaga tggcgcgttc tcatggtggg 300
ctcttcccct tggtttgtaa cgaagtntag gaggagaact tgggagccag gttctccctg 360
ccaaaaaggg ggctagatga ggtggtcggg cccgtggaca gctgagagtg ggattcatcc 420
agactcatgc aataaccctt tgattgtttc taaaaggaga ctccctcggc aagatggcag 480
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<212> DNA
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ggggcccatt cgatctggtg attgggggca gtccctgcaa tgacctctcc atcgtcaacc 240
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tgcatgatgc gcggcccaag gagggagatg atcgcccctt cttctggctc tttgagaatg 360
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<212> DNA
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cctgaagact ccgtaccctc tgccatcttg ccgagggagt ctccttttag aaaacaatca 180
aagggttatt gcatgagtct ggatgaatcc cactctcagc tgtccacggg cccgaccacc 240
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cttgcgaaaa gggttggaca tcatctcctg atttttcaat gttattcttc agtcctattt 180
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<211> 445

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tcgtggctcc agttacaaaa aaattttaat gaaaacgtta aacatanaaa tagaagtttg 180
agattttaaa aagtgtataa aaagccccac aaaacttgtc aacggttgtt ccttattcta 240
caaaatagca ccagtaagaa gagtaaaagg tgttaaaaac catttatgac agcatttctg 300
aaatgcagct tgtctgaatt cccggttctc cctaaaaacg acttctttat ggnattaaaa 360
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<223> May be any nucleic acid
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cgtggctcca gttacaaaaa aattttaatg aaaacgttaa acataaaaat agaagtttga 180
gattttaaaa agtgtataaa aagccccaca aaacttgtca acgttgttcc ttattctaca 240
aaatagcacc agtaagaaga gtaaaaggtg ttaaaaacca ttatgacagc atttctgaaa 300
tgcagcttgt ctgaattccc gttctcccta aaaacgactt cttatggaat aaaaaaggat 360
taaaaaatct ccaaagggag caccgagctt tgcagttttc cctgtccgtc tctcagatgt 420
ggggaaggta tgagaaatgt atgtctgtcc cngactgctg tcactgcctc tgagttagta 480
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tacgtgaacg acgtgaggaa catcacaaag aaaaatattg aagaatgggg cccatttgac 180

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ccaaggaggg tgatgaccgg ccgttcttct ggatgtttga gaatgttgta gccatgaagg 360
ttggcgacaa gagggacate teacggttee tggagtgtaa tecagtgatg attgatgeca 420
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<223> May be any nucleic acid
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atagcatcaa agaatgataa actcgngctg caggactgct tggaatacaa taggatagcc 180
aagttaaaga aagtacagac aataaccacc aagtcgaact cgatcaaaca ggggaaaaac 240
caacttttcc ctgttgtcat gaatggcaaa gaagatgttt ngtggtgcac tgagctcgaa 300
aggntetttg gettteetgt geactacaca gaegtgteea acatgggeeg tggtgeeege 360
cagaagctgc tgggaaggtc ctggagcgtg cctgtcatcc gacacctctt cgcccctctg 420
aaggactact ttgcatgtga atagttccag ccagggccca agcccactgg ggtgtgtggc 480
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<210> 62
<211> 573
<212> DNA
<213> Homo sapiens
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ttcatggtca cttttttgtt tatctcattt tctctgaggc tggtccttcc tgttaacgtc 180
ttggcatttg tgggaagcac aaaatgttct tgtccctcca actctgcttt tcgctccctg 240
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cootgocatt cototocogo gootgocoto tocottocat otttocoagg tacttttoto 300
teccageeet gecaetette tgeegeaeet gegeteteee etecatettt eecaggtaet 360
tttgagcctt gactccccag gtcccttcat tctgtgctca ctccatgatg tcattttgtt 420
ctccagttaa agaaagtaca gacaataacc accaagtcga actcgatcaa acaggggaaa 480
aaccaacttt teeetgttgt catgaatgge aaagaagatg ttttgtggtg caetgagete 540
gaaaggatct ttggctttcc tgtgcactac aca
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<211> 559
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<213> Homo sapiens
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gtggctccag ttacaaaaaa attttaatga aaacgttaaa cataaaaata gaagtttgag 180
attttaaaaa gtgtataaaa agccccacaa aacttgtcaa cgttgttcct tattctacaa 240
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gcagcttgtc tgaattcccg ttctccctaa aaacgacttc ttatggaata aaaaaggatt 360
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ggaaggtatg agaaatgtat gtctgtccct gactgctgtc actgcctctg agtttagtaa 480
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gtctcctgga gagggaatg
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<210> 64
<211> 391
<212> DNA
<213> Homo sapiens
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tgttacgtcg tggctccagt tacaaaaaaa attttaatga aaacgttaaa cataaaaata 180
gaagtttgag attttaaaaa gtgtataaaa agccccacaa aacttgtcaa cgttgttcct 240
tattctacaa aatagcacca gtaagaagag taaaaggtgt taaaaaccat tatgacagca 300
tttctgaaat gcagcttgtc tgaattcccg ttctccctaa aaacgacttc ttatggaata 360
aaaaaggatt aaaaaatctc caaagggagc a
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<212> DNA
<213> Homo sapiens
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gctccagtta cggaaaaatt ttaatgaaaa cgttaaacat aaaaatagaa gtttqaqatt 180
ttaaaaagtg tataaaaagc cccacaaaac ttgtcaacgt tgttccttat tctacaaaat 240
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gcttgtctga attcccgttc tccctaaaaa cgacttctta tggaataaaa aaggattaaa 360
aaateteeaa agggageace gagetttgea gtttteeetg teatetetea gatgtgggga 420
aggtatgaga aatgtatgtc tgtccctgac tgctgtcact gcctctgagt ttagtaaaaa 480
gatgagaaat gagggtagca gacttctcat ctgggga
                                                                   517
<210> 66
<211> 442
<212> DNA
<213> Homo sapiens
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ggaaggtatg agaaatgtat gt
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<211> 396
<212> DNA
<213> Homo sapiens
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aactgcaaag ctcggtgctc cctttggaga ttttttaatc ctttttttt ccataagaag 120
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tttttaacac cttttactct tcttactggt gctattttgt agaataagga acaacgttga 240
caagttttgt ggggcttttt atacactttt taaaatctca aacttctatt tttatgttta 300
acgttttcat taaaattttt ttgtaactgg agccacgacg taacaaatat ggggaaaaaa 360
ctgtgccttg tttcaacagt ttttgctaat ttttag
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ctaaaaatta gcaaaaactg ttgaaacaag gcacagtttt tcccccatat ttgttacgtc 120
gtggctccag ttacaaaaaa aattttaatg aaaacgttaa acataaaant agaagtttga 180
gattttaaaa agtgtataaa aagccccaca aaacttgtca acqttgttcc ttattctaca 240
aaatagcacc agtaagaaga gtaaaaggtg ttaaaaacca ttatgac
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<210> 69
<211> 356
<212> DNA
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<223> May be any nucleic acid
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accacctgct gaattactca cgccccaagg agggtgatga ccggccgttc ttctggatgt 180
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gtaatccagt gatgattgat gccatcaaag tttctgctgc tcacagggcc cgatacttct 300
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<222> (408)..(408)
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gtcgtggctc cagttacaaa aaaaatttta atgaaaacgt taaacataaa aatagaagtt 180
tgagatttta aaaagtgtat aaaaagcccc acaaaacttg tcaacgttgt tccttattct 240
acaaaatagc accagtaaga agagtaaaaag gtgttaaaaa ccattatgac agcatttctg 300
aaatgcagct tgtctgaatt cccgttctcc ctaaaaacga cttcttatgg aataaaaaag 360
gattaaaaaa tctccaaagg gagcaccgag ctttgcagtt ttccctgn
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aaaaaaaatt ttaatgaaaa cgttaaacat aaaaatagaa gtttgagatt ttaaaaagtg 180
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gtatggggaa atggtatggt ctggtccctg gactggctgg tcactgcctc tggggtttng 480
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accggncgtt cttctggatg tttgagaatg ttgtagncat gaaggttggn gacaagaggg 180
acateteacg gtteetggag tgtaateeag tgatgattga tgceateaaa gtttetgetg 240
ctcacagggc ccgatacttc tggggcaacc tacccgggat gaacaggatc tttggctttc 300
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tggggaaggt nctggggagc gttgccttgt tcatcccgac acctntttcg gnccctattg 420
gaagggattn atttttgcca tgt
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<211> 273
<212> DNA
<213> Homo sapiens
<400> 74
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ctggattagt tctgatccca ccacaaggag ccctcgaatt ggctaaagtg agaaactggg 120
cctgaagact ccgtaccctc tgccatcttg ccgagggagt ctccttttag aaaacaatca 180
aagggttatt gcatgagtct ggatgaatcc cactctcagc tgtccacggg cccgaccacc 240
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aaaaaaaatt ttaatgaaaa cgttaaacat aaaaatagaa gtttgagatt ttaaaaagtg 180
tataaaango cccacaaaac ttgtcaacgt tgttccttat tctacaaaat agcaccagta 240
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Lys Asn Ala Thr Lys Ala Ala Asp Asp Glu Pro Glu Tyr Glu Asp Gly 50 55 60

Arg Gly Phe Gly Ile Gly Glu Leu Val Trp Gly Lys Leu Arg Gly Phe 65 70 75 80

Ser Trp Trp Pro Gly Arg Ile Val Ser Trp Trp Met Thr Gly Arg Ser

/ 85 90 95

Arg Ala Ala Glu Gly Thr Arg Trp Val Met Trp Phe Gly Asp Gly Lys 100 105 110

Phe Ser Val Val Cys Val Glu Lys Leu Met Pro Leu Ser Ser Phe Cys 115 120 125

Ser Ala Phe His Gln Ala Thr Tyr Asn Lys Gln Pro Met Tyr Arg Lys 130 135 140

Ala Ile Tyr Glu Val Leu Gln Val Ala Ser Ser Arg Ala Gly Lys Leu 145 150 155 160

Phe Pro Ala Cys His Asp Ser Asp Glu Ser Asp Ser Gly Lys Ala Val 165 170 175

Glu Val Gln Asn Lys Gln Met Ile Glu Trp Ala Leu Gly Gly Phe Gln 180 185 190

Pro Ser Gly Pro Lys Gly Leu Glu Pro Pro Glu Glu Glu Lys Asn Pro 195 200 205

Tyr Lys Glu Val Tyr Thr Asp Met Trp Val Glu Pro Glu Ala Ala Ala 210 215 220

Tyr Ala Pro Pro Pro Pro Ala Lys Lys Pro Arg Lys Ser Thr Thr Glu 225 230 235 240

Lys Pro Lys Val Lys Glu Ile Ile Asp Glu Arg Thr Arg Glu Arg L'eu 245 250 255

Val Tyr Glu Val Arg Gln Lys Cys Arg Asn Ile Glu Asp Ile Cys Ile 260 265 270

Ser Cys Gly Ser Leu Asn Val Thr Leu Glu His Pro Leu Phe Ile Gly 275 280 285

Gly Met Cys Gln Asn Cys Lys Asn Cys Phe Leu Glu Cys Ala Tyr Gln 290 295 300

Tyr Asp Asp Gly Tyr Gln Ser Tyr Cys Thr Ile Cys Cys Gly Gly 305 310 315 320

Arg Glu Val Leu Met Cys Gly Asn Asn Cys Cys Arg Cys Phe Cys 325 330 335

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Ile Lys Glu Asp Pro Trp Asn Cys Tyr Met Cys Gly His Lys Gly Thr 355 360 365

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Gln Val Asp Arg Tyr Ile Ala Ser Glu Val Cys Glu Asp Ser Ile Thr 435 440 445

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Arg Ser Val Thr Gln Lys His Ile Gln Glu Trp Gly Pro Phe Asp Leu 465 470 475 480

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515 520 525

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Asp Ile Ser Arg Phe Leu Glu Ser Asn Pro Val Met Ile Asp Ala Lys 545 550 555 560

Glu Val Ser Ala Ala His Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro 565 570 575

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Gln Glu Cys Leu Glu His Gly Arg Ile Ala Lys Phe Ser Lys Val Arg 595 600 605

Thr Ile Thr Thr Arg Ser Asn Ser Ile Lys Gln Gly Lys Asp Gln His 610 620

Phe Pro Val Phe Met Asn Glu Lys Glu Asp Ile Leu Trp Cys Thr Glu 625 630 635 640

Met Glu Arg Val Phe Gly Phe Pro Val His Tyr Thr Asp Val Ser Asn 645 650 655

Met Ser Arg Leu Ala Arg Gln Arg Leu Leu Gly Arg Ser Trp Ser Val 660 665 670

Pro Val Ile Arg His Leu Phe Ala Pro Leu Lys Glu Tyr Phe Ala Cys 675 680 685

Val

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Lys Asn Ala Thr Lys Ala Gly Asp Asp Glu Pro Glu Tyr Glu Asp Gly 50 55 60

Arg Gly Phe Gly Ile Gly Glu Leu Val Trp Gly Lys Leu Arg Gly Phe 65 70 75 80

Ser Trp Trp Pro Gly Arg Ile Val Ser Trp Trp Met Thr Gly Arg Ser 85 90 95

Arg Ala Ala Glu Gly Thr Arg Trp Val Met Trp Phe Gly Asp Gly Lys 100 105 110

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Phe Pro Val Cys His Asp Ser Asp Glu Ser Asp Thr Ala Lys Ala Val 165 170 175

Glu Val Gln Asn Lys Pro Met Ile Glu Trp Ala Leu Gly Gly Phe Gln 180 185 190

Pro Ser Gly Pro Lys Gly Leu Glu Pro Pro Glu Glu Glu Lys Asn Pro 195 200 205

Tyr Lys Glu Val Tyr Thr Asp Met Trp Val Glu Pro Glu Ala Ala Ala 210 215 220

Tyr Ala Pro Pro Pro Pro Ala Lys Lys Pro Arg Lys Ser Thr Ala Glu 225 230 235 240

Lys Pro Lys Val Lys Glu Ile Ile Asp Glu Arg Thr Arg Glu Arg Leu 245 250 255

Val Tyr Glu Val Arg Gln Lys Cys Arg Asn Ile Glu Asp Ile Cys Ile 260 265 270

Ser Cys Gly Ser Leu Asn Val Thr Leu Glu His Pro Leu Phe Val Gly 275 280 285

Gly Met Cys Gln Asn Cys Lys Asn Cys Phe Leu Glu Cys Ala Tyr Gln 290 295 300

Tyr Asp Asp Asp Gly Tyr Gln Ser Tyr Cys Thr Ile Cys Cys Gly Gly 305 310 315

Arg Glu Val Leu Met Cys Gly Asn Asn Cys Cys Arg Cys Phe Cys 325 330 335

Val Glu Cys Val Asp Leu Leu Val Gly Pro Gly Ala Ala Gln Ala Ala 340 345 350

Ile Lys Glu Asp Pro Trp Asn Cys Tyr Met Cys Gly His Lys Gly Thr 355 360 365

Tyr Gly Leu Leu Arg Arg Glu Asp Trp Pro Ser Arg Leu Gln Met 370 375 380

Phe Phe Ala Asn Asn His Asp Gln Glu Phe Asp Pro Pro Lys Val Tyr 385 390 395 400

Pro Pro Val Pro Ala Glu Lys Arg Lys Pro Ile Arg Val Leu Ser Leu 405 410 415

Phe Asp Gly Ile Ala Thr Gly Leu Leu Val Leu Lys Asp Leu Gly Ile 420 425 430

Gln Val Asp Arg Tyr Ile Ala Ser Glu Val Cys Glu Asp Ser Ile Thr 435 440 445

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Val Ile Gly Gly Ser Pro Cys Asn Asp Leu Ser Ile Val Asn Pro Ala 485 490 495

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Arg Leu Leu His Asp Ala Arg Pro Lys Glu Gly Asp Asp Arg Pro Phe

515 520 525

Phe Trp Leu Phe Glu Asn Val Val Ala Met Gly Val Ser Asp Lys Arg 530 540

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Thr Ile Thr Thr Arg Ser Asn Ser Ile Lys Gln Gly Lys Asp Gln His 610 620

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